

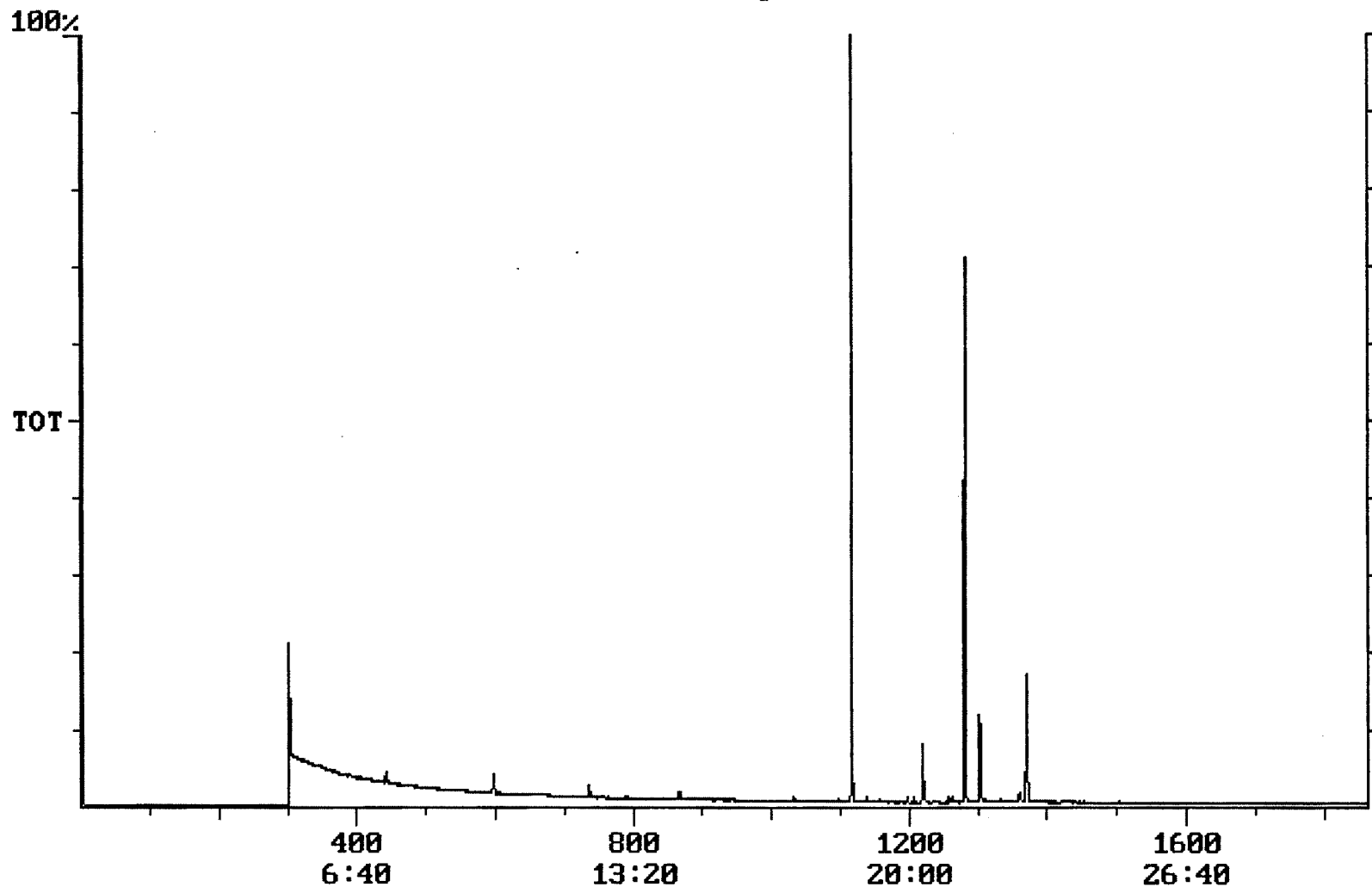
User: Bohlk, Ted
 Westchester Dept. of Labs & Research
 Date: 10/10/2001 Time: 08:59:44

Sample: AD23082
 Analysis: \$LP525 Organic Extractables
 Units: ug
 Started: 10/4/01 08:58 Ended: 10/4/01 08:58 Analyst: TB
 Page: 1

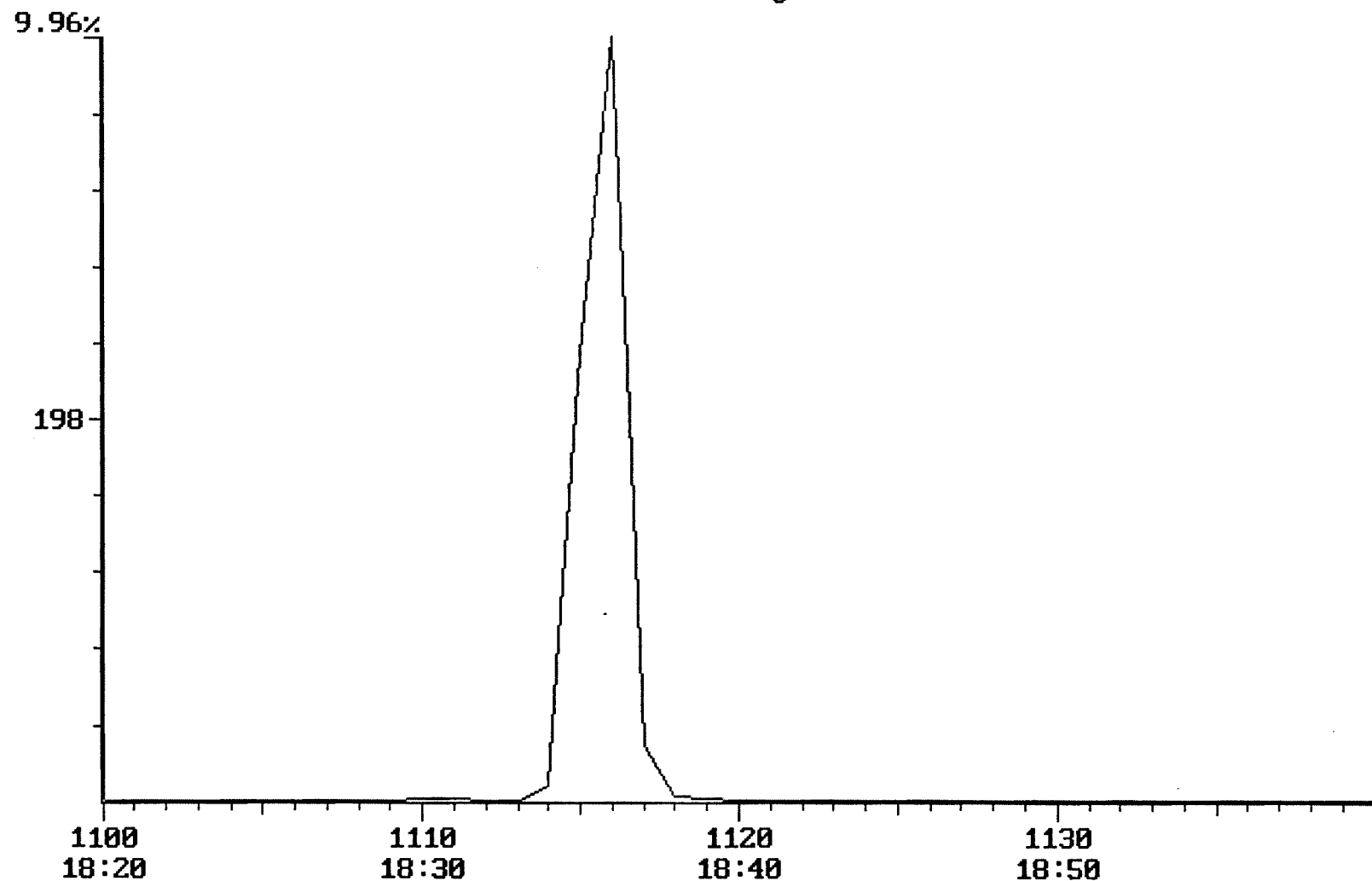
	Component Name	Viol	Result
1	DFTPP		PASSED
2	ENDRIN *		37
3	DDT		2.2

→ flag

Chromatogram Plot File: E:\DFTA1004 Date: Oct-04-1991 17:05:32
Comment: BOHLK; METHOD 525; INST 204; 10/04/01; DFIPP
Scan No: 1 Retention Time: 0:01 RIC: 0 Mass Range: 0 - 0
Plotted: 1 to 1859 Range: 1 to 1859 100% = 4498629



Chromatogram Plot File: E:\DFTA1004 Date: Oct-04-1991 17:05:32
Comment: BOHLK; METHOD 525; INST 204; 10/04/01; DFTPP
Scan No: 1100 Retention Time: 18:20 RIC: 30265 Mass Range: 47 - 86
Plotted: 1100 to 1140 Range: 1 to 1859 100% = 4498629



Mass List

File: E:\DFTA1004 Date: Oct-04-1991 17:05:32

Comment: BOHLK; METHOD 525; INST 204; 10/04/01; DFTPP

Scan No: 1116 Retention Time: 18:36 RIC: 4498629 Mass Range: 47 - 444

Peaks: 159 Mass of Base Peak: 442 Int: 758762 Threshold: 0 %

Mass	Intensity	% Base	Mass	Intensity	% Base	Mass	Intensity	% Base
47	3,092	0.41	84	3,865	0.51	123	6,185	0.82
49	14,948	1.97	85	5,154	0.68	124	4,123	0.54
50	100,773	13.28	86	10,567	1.39	127	138,402	18.24
51	283,247	37.33	87	2,835	0.37	128	13,917	1.83
52	11,340	1.49	91	3,608	0.48	129	63,402	8.36
55	5,154	0.68	92	3,865	0.51	130	4,381	0.58
56	8,505	1.12	93	35,309	4.65	135	6,958	0.92
57	11,597	1.53	94	2,061	0.27	136	2,835	0.37
61	4,896	0.65	96	1,804	0.24	141	10,824	1.43
62	5,412	0.71	98	33,247	4.38	142	3,350	0.44
63	7,216	0.95	99	14,432	1.90	143	3,092	0.41
68	3,608	0.48	100	2,319	0.31	146	2,577	0.34
69	212,886	28.06	101	4,639	0.61	147	4,896	0.65
71	1,288	0.17	103	5,670	0.75	148	12,886	1.70
73	4,381	0.58	104	6,701	0.88	149	3,865	0.51
74	35,824	4.72	105	7,731	1.02	153	4,639	0.61
75	27,577	3.63	106	2,061	0.27	154	2,835	0.37
76	10,051	1.32	107	67,783	8.93	155	5,670	0.75
77	214,432	28.26	108	7,731	1.02	156	9,020	1.19
78	12,628	1.66	110	113,659	14.98	160	3,350	0.44
79	28,865	3.80	111	15,206	2.00	161	5,670	0.75
80	14,432	1.90	112	2,061	0.27	165	2,577	0.34
81	15,721	2.07	117	87,113	11.48	166	4,639	0.61
82	4,896	0.65	118	5,154	0.68	167	23,969	3.16
83	3,092	0.41	122	3,608	0.48	168	7,989	1.05

Mass List

File: E:\DFTA1004 Date: Oct-04-1991 17:05:32

Comment: BOHLK; METHOD 525; INST 204; 10/04/01; DFTPP

Scan No: 1116 Retention Time: 18:36 RIC: 4498629 Mass Range: 47 - 444

Peaks: 159 Mass of Base Peak: 442 Int: 758762 Threshold: 0 %

Mass	Intensity	% Base	Mass	Intensity	% Base	Mass	Intensity	% Base
168	7,989	1.05	207	12,371	1.63	275	131,443	17.32
172	1,804	0.24	208	3,092	0.41	276	17,010	2.24
173	3,092	0.41	215	1,804	0.24	277	12,628	1.66
174	2,835	0.37	217	22,164	2.92	278	3,092	0.41
175	5,154	0.68	218	3,608	0.48	285	2,061	0.27
176	2,319	0.31	221	7,731	1.02	293	3,865	0.51
177	4,381	0.58	223	5,154	0.68	296	28,350	3.74
179	12,113	1.60	224	39,690	5.23	297	5,154	0.68
180	7,216	0.95	225	11,855	1.56	303	3,865	0.51
181	4,381	0.58	227	20,876	2.75	315	4,123	0.54
185	8,247	1.09	228	3,092	0.41	316	3,350	0.44
186	54,639	7.20	229	3,608	0.48	323	14,175	1.87
187	17,010	2.24	235	2,577	0.34	324	3,350	0.44
188	2,061	0.27	244	28,350	3.74	327	3,350	0.44
189	3,608	0.48	245	3,608	0.48	334	9,793	1.29
192	4,381	0.58	246	7,989	1.05	346	2,577	0.34
193	2,061	0.27	249	2,835	0.37	352	3,608	0.48
196	8,505	1.12	255	257,731	33.97	353	3,350	0.44
197	4,123	0.54	256	37,371	4.93	354	7,216	0.95
198	448,195	59.07	257	2,577	0.34	365	34,793	4.59
199	22,938	3.02	258	16,237	2.14	366	5,927	0.78
203	3,608	0.48	265	5,154	0.68	371	2,319	0.31
204	11,340	1.49	266	1,546	0.20	372	8,762	1.15
205	20,103	2.65	273	9,536	1.26	373	3,608	0.48
206	73,195	9.65	274	29,123	3.84	390	1,546	0.20

Mass List

File: E:\DFTA1004 Date: Oct-04-1991 17:05:32

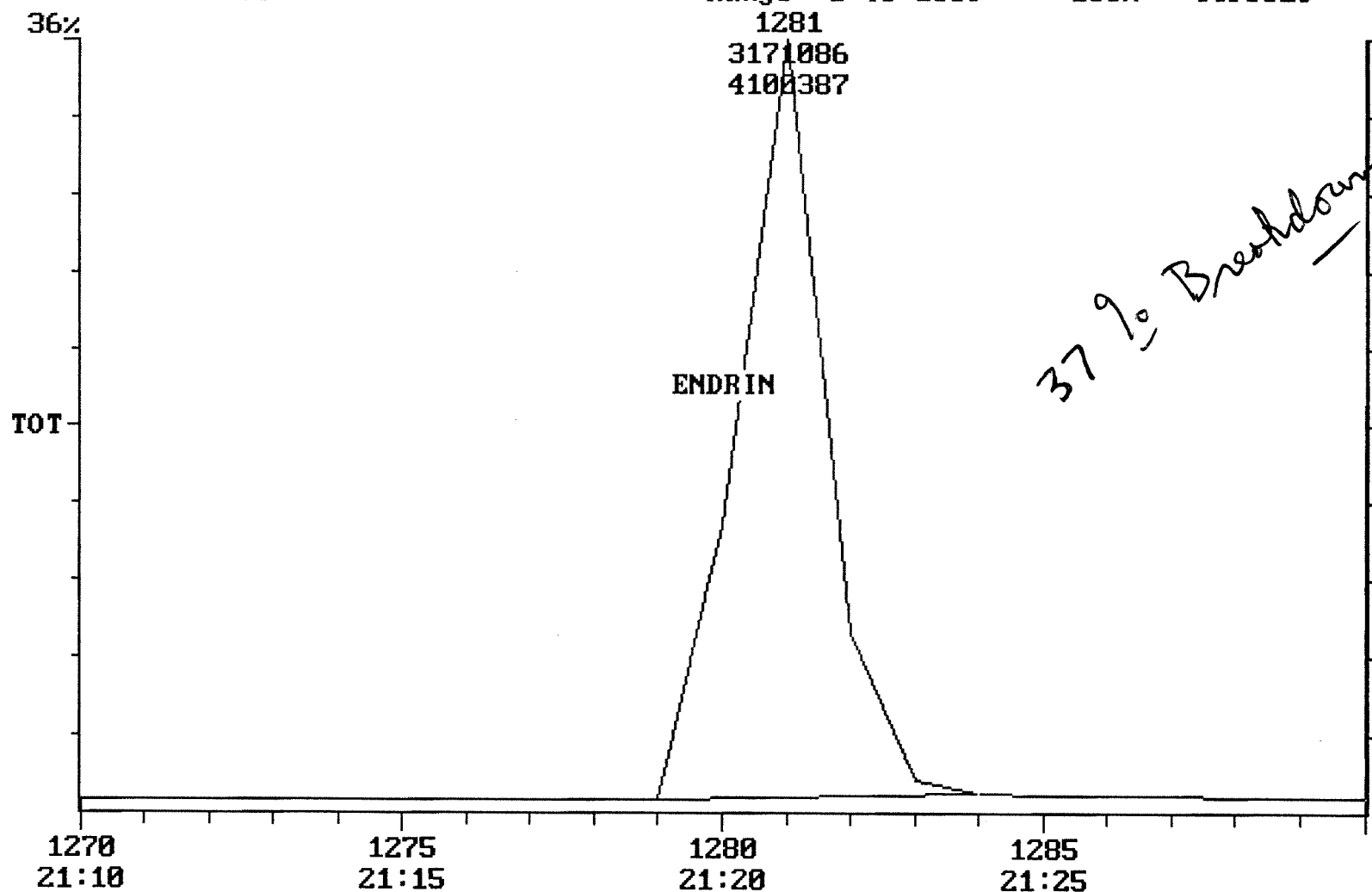
Comment: BOHLK; METHOD 525; INST 204; 10/04/01; DFTPP

Scan No: 1116 Retention Time: 18:36 RIC: 4498629 Mass Range: 47 - 444

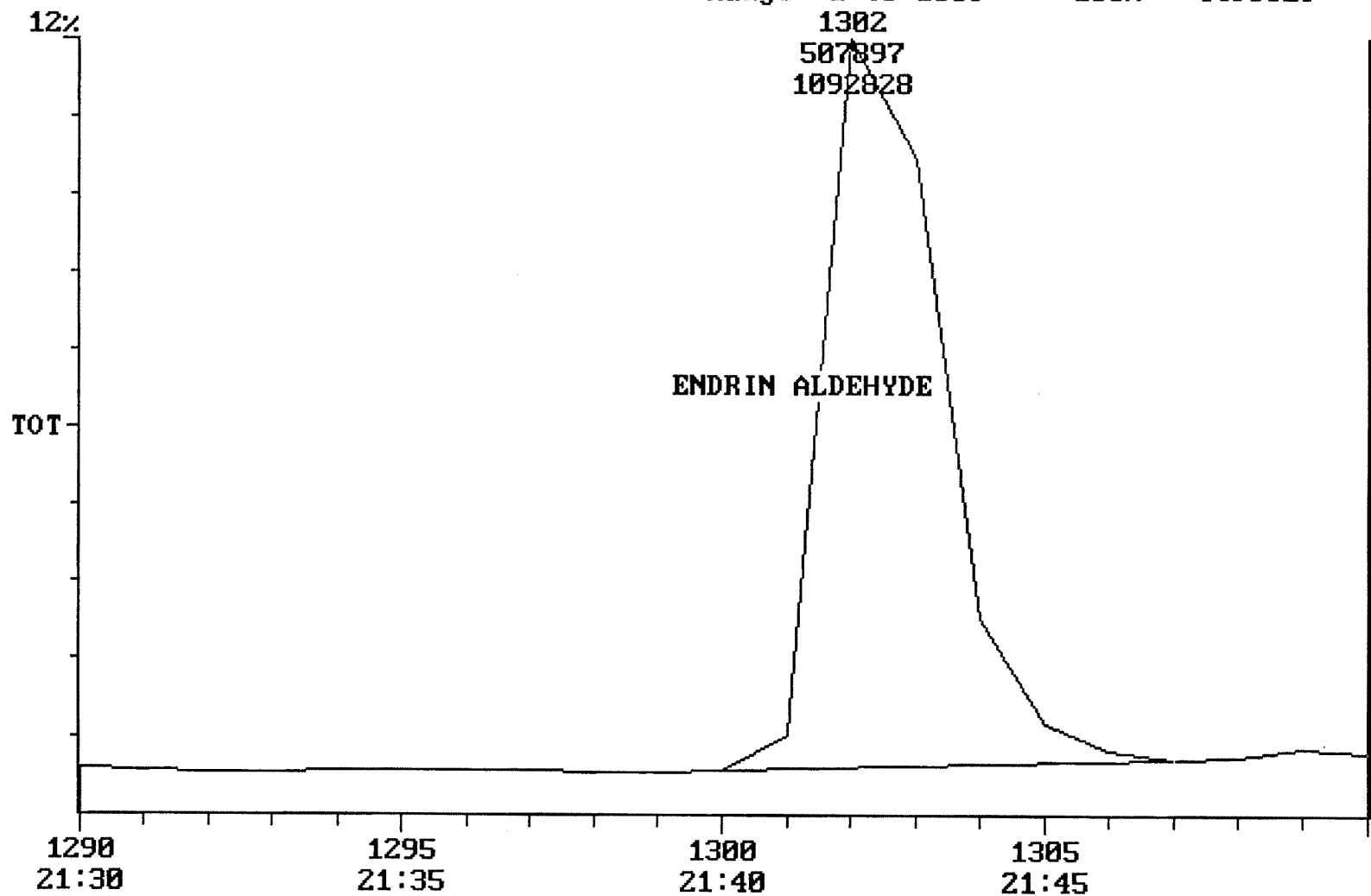
Peaks: 159 Mass of Base Peak: 442 Int: 758762 Threshold: 0 %

Mass	Intensity	% Base	Mass	Intensity	% Base	Mass	Intensity	% Base
185	8,247	1.09	228	3,092	0.41	316	3,350	0.44
186	54,639	7.20	229	3,608	0.48	323	14,175	1.87
187	17,010	2.24	235	2,577	0.34	324	3,350	0.44
188	2,061	0.27	244	28,350	3.74	327	3,350	0.44
189	3,608	0.48	245	3,608	0.48	334	9,793	1.29
192	4,381	0.58	246	7,989	1.05	346	2,577	0.34
193	2,061	0.27	249	2,835	0.37	352	3,608	0.48
196	8,505	1.12	255	257,731	33.97	353	3,350	0.44
197	4,123	0.54	256	37,371	4.93	354	7,216	0.95
198	448,195	59.07	257	2,577	0.34	365	34,793	4.59
199	22,938	3.02	258	16,237	2.14	366	5,927	0.78
203	3,608	0.48	265	5,154	0.68	371	2,319	0.31
204	11,340	1.49	266	1,546	0.20	372	8,762	1.15
205	20,103	2.65	273	9,536	1.26	373	3,608	0.48
206	73,195	9.65	274	29,123	3.84	390	1,546	0.20
207	12,371	1.63	275	131,443	17.32	402	3,608	0.48
208	3,092	0.41	276	17,010	2.24	403	3,865	0.51
215	1,804	0.24	277	12,628	1.66	421	4,896	0.65
217	22,164	2.92	278	3,092	0.41	422	5,927	0.78
218	3,608	0.48	285	2,061	0.27	423	42,268	5.57
221	7,731	1.02	293	3,865	0.51	424	6,185	0.82
223	5,154	0.68	296	28,350	3.74	441	103,092	13.59
224	39,690	5.23	297	5,154	0.68	442	758,762	100.00
225	11,855	1.56	303	3,865	0.51	443	124,484	16.41
227	20,876	2.75	315	4,123	0.54	444	13,144	1.73

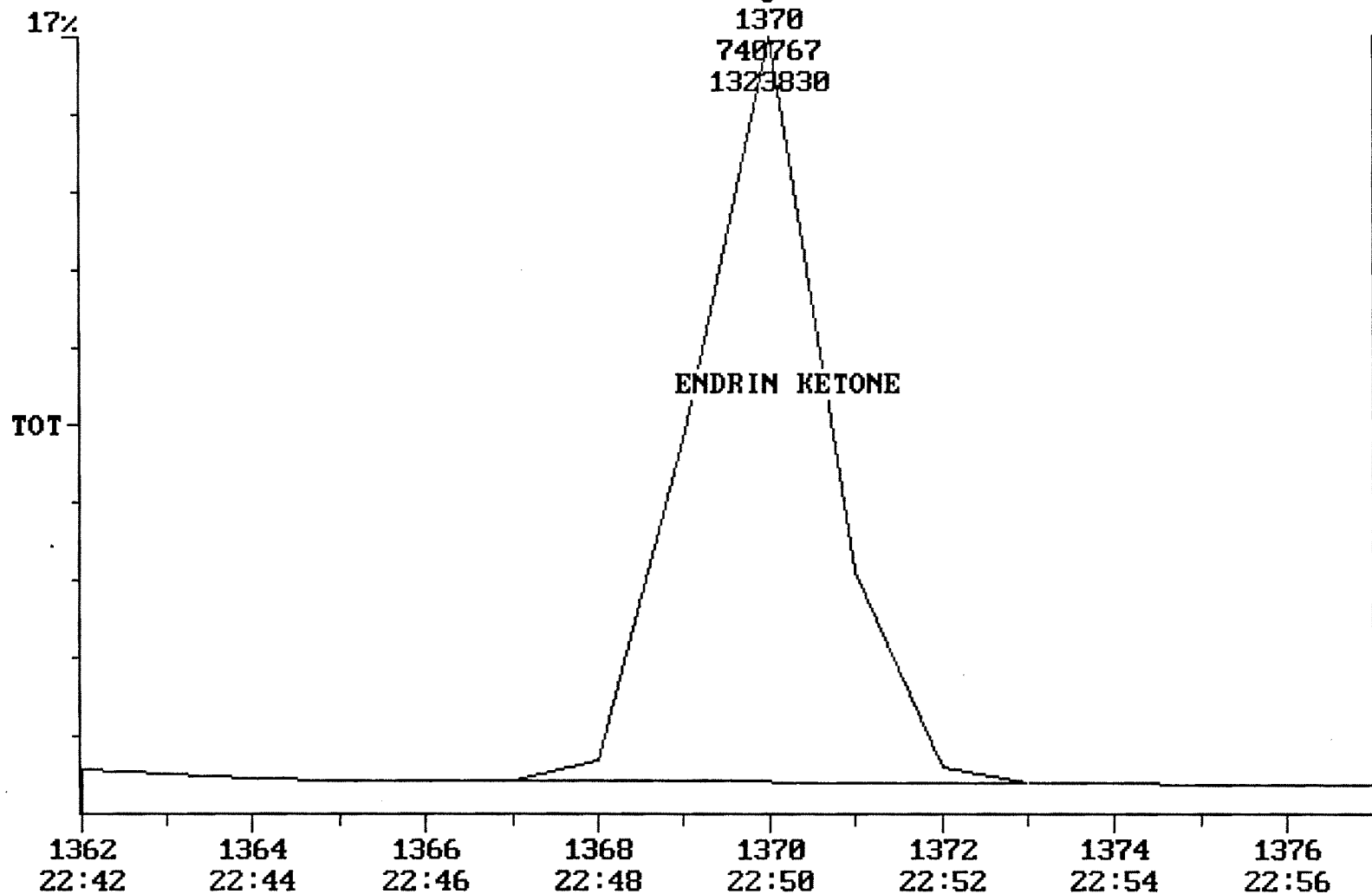
Chromatogram Plot File: E:\DFTA1004 Date: Oct-04-1991 17:05:32
 Comment: BOHLK; METHOD 525; INST 204; 10/04/01; DFTPP
 Scan No: 1290 Retention Time: 21:30 RIC: 31834 Mass Range: 45 - 281
 Plotted: 1270 to 1290 Range: 1 to 1859 100% = 4498629



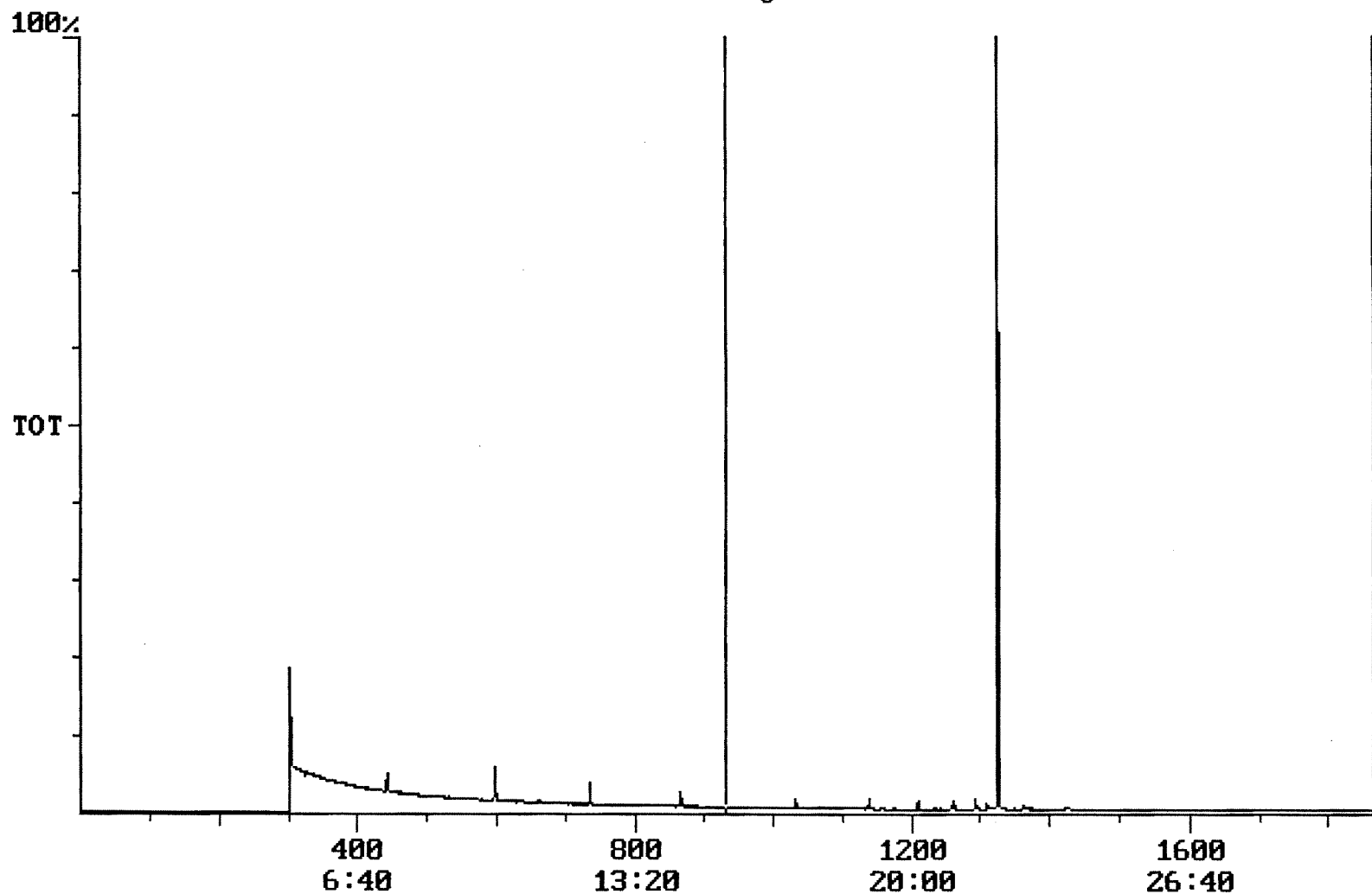
Chromatogram Plot File: E:\DFTA1004 Date: Oct-04-1991 17:05:32
Comment: BOHLK; METHOD 525; INST 204; 10/04/01; DFTPP
Scan No: 1310 Retention Time: 21:50 RIC: 42070 Mass Range: 45 - 355
Plotted: 1290 to 1310 Range: 1 to 1859 100% = 4498629



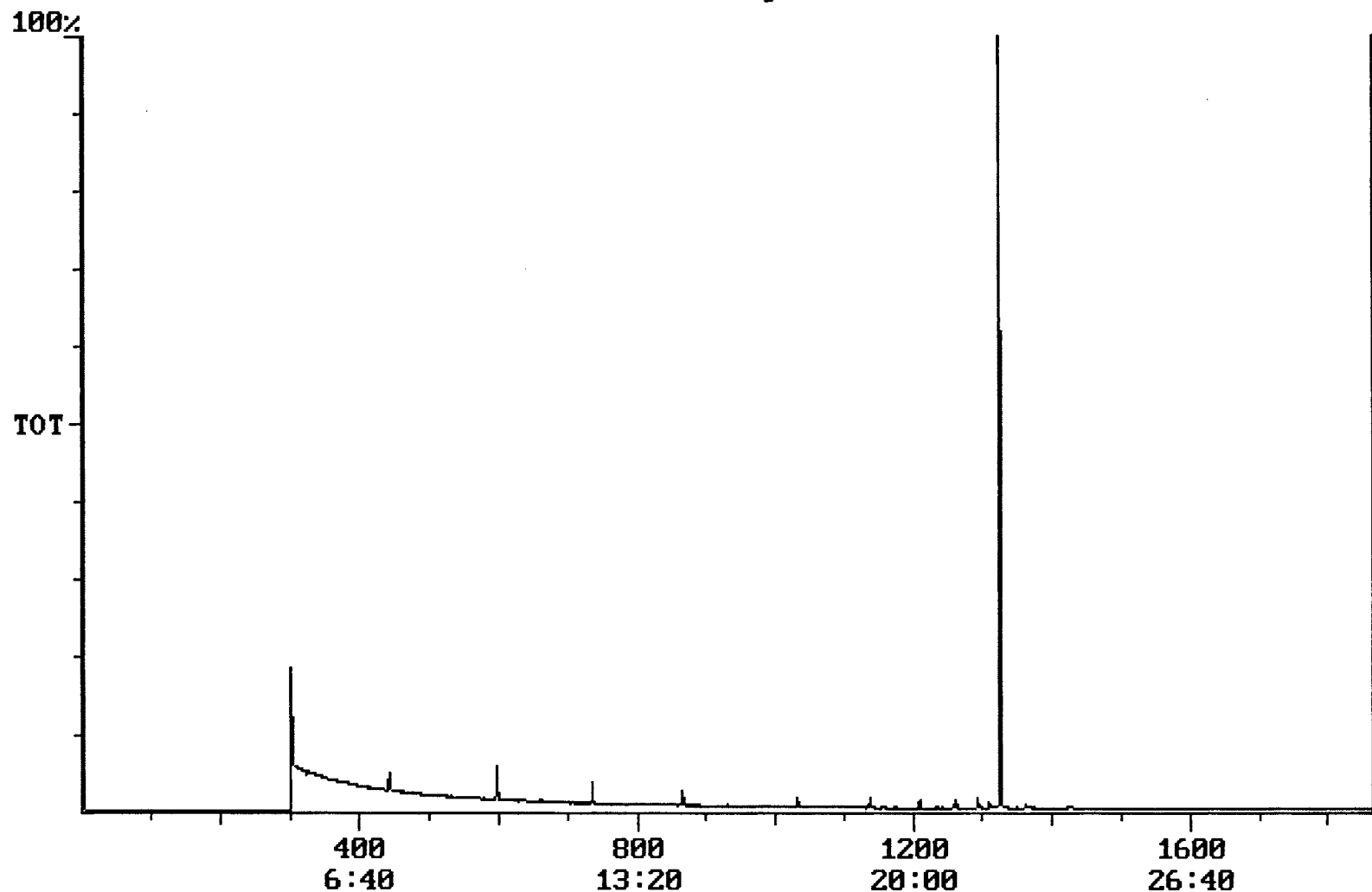
Chromatogram Plot File: E:\DFTA1004 Date: Oct-04-1991 17:05:32
Comment: BOHLK; METHOD 525; INST 204; 10/04/01; DFTPP
Scan No: 1377 Retention Time: 22:57 RIC: 27539 Mass Range: 45 - 281
Plotted: 1362 to 1377 Range: 1 to 1859 100% = 4498629



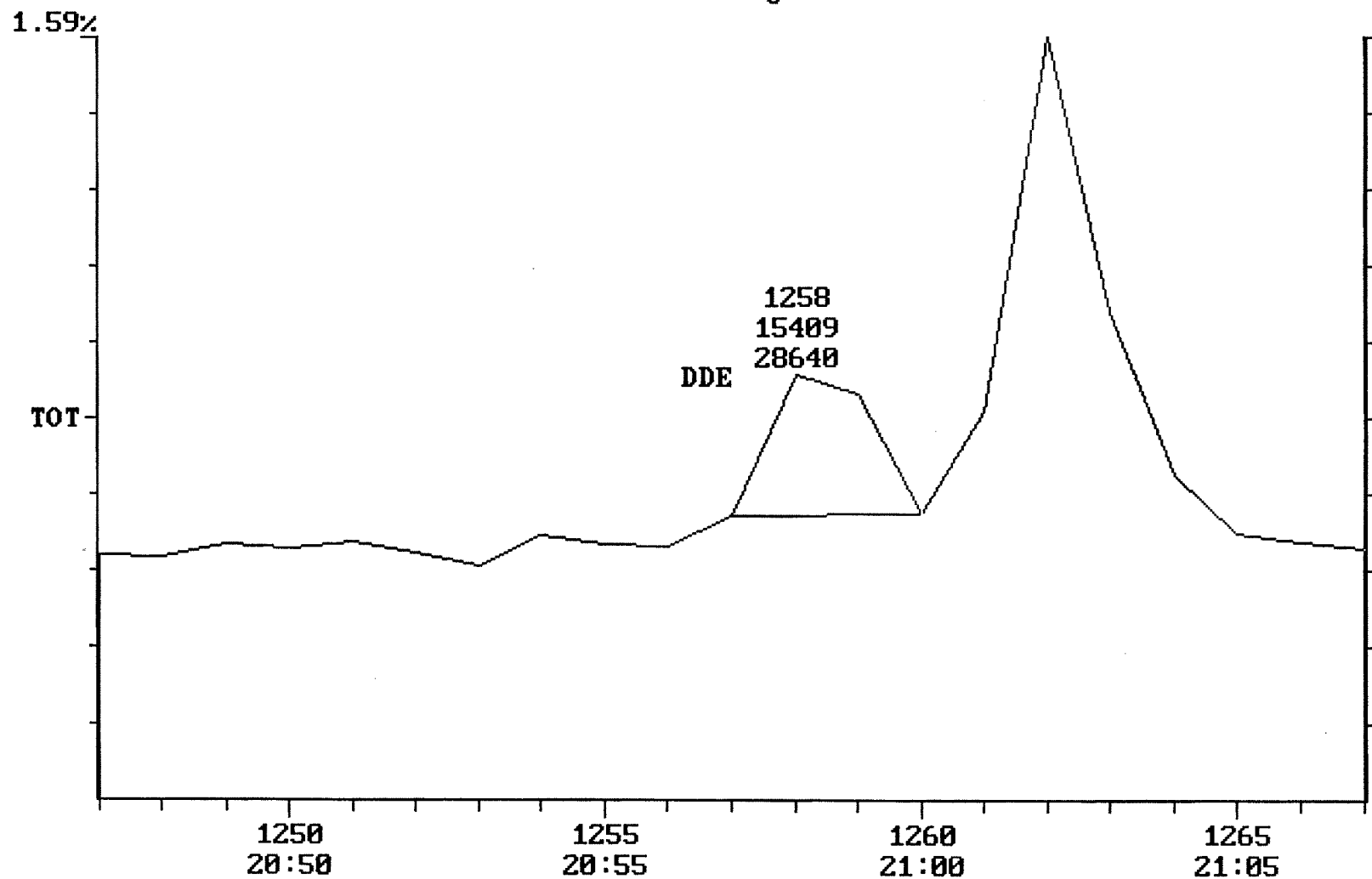
Chromatogram Plot File: E:\DDTA1004 Date: Oct-04-1991 17:42:08
Comment: BOHLK; METHOD 525; INST 204; 10/04/01; DDT
Scan No: 930 Retention Time: 15:30 RIC: 44160 Mass Range: 47 - 86
Plotted: 1 to 1860 Range: 1 to 1860 100% = 5248161



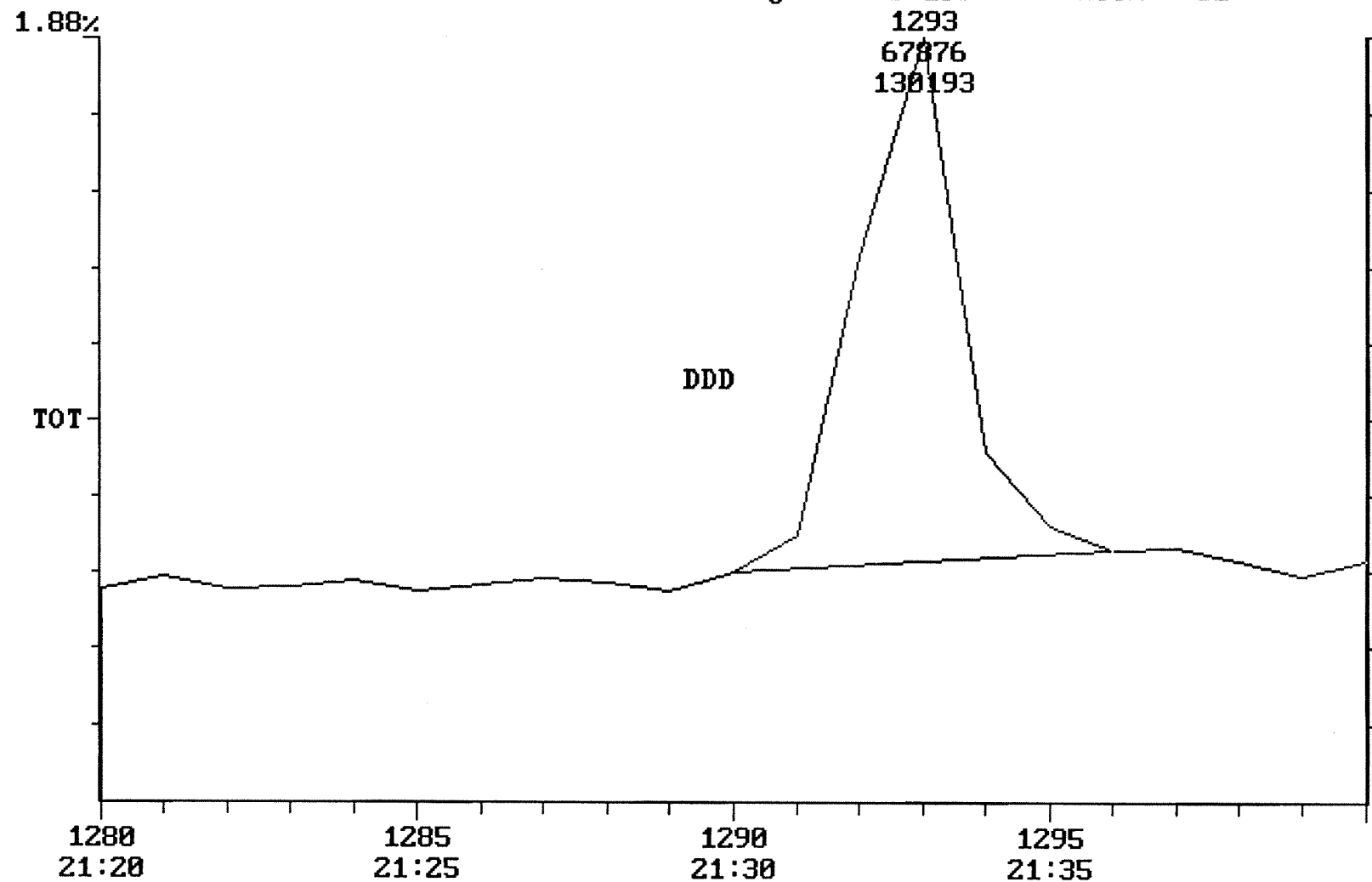
Chromatogram Plot File: E:\DDTA1004 Date: Oct-04-1991 17:42:08
Comment: BOHLK; METHOD 525; INST 204; 10/04/01; DDT
Scan No: 1 Retention Time: 0:01 RIC: 0 Mass Range: 0 - 0
Plotted: 1 to 1860 Range: 1 to 1860 100% = 5248161



Chromatogram Plot File: E:\DDTA1004 Date: Oct-04-1991 17:42:08
Comment: BOHLK; METHOD 525; INST 204; 10/04/01; DDT
Scan No: 1267 Retention Time: 21:07 RIC: 27388 Mass Range: 47 - 207
Plotted: 1247 to 1267 Range: 1 to 1860 100% = 5248161



Chromatogram Plot File: E:\DDTA1004 Date: Oct-04-1991 17:42:08
 Comment: BOHLK; METHOD 525; INST 204; 10/04/01; DDT
 Scan No: 1300 Retention Time: 21:40 RIC: 30956 Mass Range: 45 - 235
 Plotted: 1280 to 1300 Range: 1 to 1860 100% = 5248161



Chromatogram Plot File: E:\DDTA1004 Date: Oct-04-1991 17:42:08
 Comment: BOHLK; METHOD 525; INST 204; 10/04/01; DDT
 Scan No: 1333 Retention Time: 22:13 RIC: 39767 Mass Range: 45 - 237
 Plotted: 1313 to 1333 Range: 1 to 1860 100% = 5248161

